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<222> (8)
<223> Xaa is Ala, Gln, Gly, Lys or Thr
<220>
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<222> (9)
<223> Xaa is Arg, Asn, Asp, Glu or Gly
<220>
<221> VARIANT '
<222> (10)
<223> Xaa is Gln, Leu or Gly
<220>
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<222> (11)
<223> Xaa is Ala, Trp or Tyr
<220>
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<222> (12)
<223> Xaa is Ala, Gly, His, Phe, Thr or Val
<220>
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<222> (14)
<223> Xaa is Asn, Gln, Phe, Ser or Val
<220>
<221> VARIANT
<222> (15)
<223> Xaa is Arg, Leu, Pro or Ser
<220>
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<222> (16)
<223> Xaa is Leu, Ser, Trp or Tyr
Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa
<210> 2
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: family of
     preferred CEA binding moieties
<220>
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<222> (1)
<223> Xaa is Asn or Asp
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<220>
<221> VARIANT
<222> (6)
<223> Xaa is Phe, Met, Leu or Asn
<220>
<221> VARIANT
<222> (7)
<223> Xaa is Asp, Gly, Ile, Lys, Phe or Thr
<220>
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<222> (9)
<223> Xaa is Arg, Asn, Asp, Glu, Gly or Trp
<220>
<221> VARIANT
<222> (12)
<223> Xaa is Ala, Gly, His, Phe, Thr, Tyr or Val
<220>
<221> VARIANT
<222> (15)
<223> Xaa is Arg, Leu, Pro or Ser
<220>
<221> VARIANT
<222> (16)
<223> Xaa is Leu, Ser, Trp or Tyr
Xaa Trp Val Cys Glu Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Xaa Xaa
                                      10
<210> 3
<211> 10
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: CEA binding
      loop
<220>
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<223> Xaa is Asn, Glu or Met
<220>
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<222> (3)
<223> Xaa is Asn, Leu, Met or Phe
<220>
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<221> VARIANT

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<222> (4)
 <223> Xaa is Asp, Gly, Ile, Lys, Phe or Thr
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 <222> (5)
 <223> Xaa is Ala, Gln, Gly, Lys or Thr
 <220>
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 <222> (6)
<223> Xaa is Arg, Asn, Asp, Glu or Gly
 <220>
 <221> VARIANT
 <222> (7)
 <223> Xaa is Gln, Gly or Leu
 <220>
 <221> VARIANT
 <222> (8)
 <223> Xaa is Ala, Trp or Tyr
 <220>
 <221> VARIANT
 <222> (9)
 <223> Xaa is Ala, Gly, His, Phe, Thr or Val
 Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys
                   5
 <210> 4
 <211> 16
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: CEA binding
       polypeptide
 <400> 4
 Asn Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asn Ser Tyr
   1
                                       10
 <210> 5
 <211> 16
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: CEA binding
       polypeptide
 <400> 5
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Asp Trp Val Cys Glu Asn Lys Lys Asp Gln Trp Thr Cys Asn Leu Leu
                                       10
<210> 6
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 6
Asn Trp Asp Cys Met Phe Gly Ala Glu Gly Trp Ala Cys Ser Pro Trp
                                      10
<210> 7
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 7
Asp Trp Val Cys Glu Lys Thr Thr Gly Gly Tyr Val Cys Gln Pro Leu
                                      10
<210> 8
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
Asn Trp Phe Cys Glu Met Ile Gly Arg Gln Trp Gly Cys Val Pro Ser
                  5
                                                           15
<210> 9
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 9
Asp Trp Val Cys Asn Phe Asp Gln Gly Leu Ala His Cys Phe Pro Ser
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1

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<210> 10
<211> 12
<212> PRT
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<220>
<223> Description of Artificial Sequence: parental
      domain for design of microprotein display library
<220>
<221> VARIANT
<222> (1)..(12)
<223> amino acid positions 4 and 9 are invariant Cys;
      all other positions Xaa are varied but not Cys, to
      provide a library of 2x10(8) different peptides
      based on the template sequence
<400> 10
Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa
<210> 11
<211> 11
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: parental
      domain for design of microprotein display library
<220>
<221> VARIANT
<222> (1)..(11)
<223> amino acid positions 3 and 9 are invariant Cys;
      all other positions Xaa are varied but not Cys, to
      provide a library of 1x10(9) different peptides
      based on the template sequence
Xaa Xaa Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa
                  5
<210> 12
<211> 12
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: parental
      domain for design of microprotein display library
<220>
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<221> VARIANT
<222> (1)..(12)
<223> amino acid positions 3 and 10 are invariant Cys;
      all other positions Xaa are varied but not Cys, to
      provide a library of 1x10(9) different peptides
      based on the template sequence
<400> 12
Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
                  5
<210> 13
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: parental
      domain for design of microprotein display library
<220>
<221> VARIANT
<222> (1)..(16)
<223> amino acid positions 4 and 13 are invariant Cys;
      all other positions Xaa are varied but not Cys, to
      provide a library of 2.5x10(8) different peptides
      based on the template sequence
<400> 13
Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa
<210> 14
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: variable
      sublibrary sequence used in designing focused
      secondary library
<220>
<221> VARIANT
<222> (1)..(3)
<223> Xaa is any amino acid except Cys
<220>
<221> VARIANT
<222> (5)..(6)
<223> Xaa is any amino acid except Cys
<400> 14
Xaa Xaa Cys Xaa Xaa Lys Lys Asp Gln Trp Thr Cys Asn Leu Leu
```

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<210> 15
 <211> 16
<212> PRT
 <213> Artificial Sequence
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 <223> Description of Artificial Sequence: variable
       sublibrary sequence used in designing focused
       secondary library
 <220>
 <221> VARIANT
 <222> (5)..(9)
 <223> Xaa is any amino acid except Cys
 Asp Trp Val Cys Xaa Xaa Xaa Xaa Gln Trp Thr Cys Asn Leu Leu
   1
                                       10
 <210> 16
 <211> 16
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: variable
       sublibrary sequence used in designing focused
       secondary library
 <220>
 <221> VARIANT
 <222> (8)..(12)
 <223> Xaa is any amino acid except Cys
 Asp Trp Val Cys Glu Asn Lys Xaa Xaa Xaa Xaa Cys Asn Leu Leu
   1
                   5
                                       10
                                                            15
 <210> 17
 <211> 16
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: variable
       sublibrary sequence used in designing focused
       secondary library
 <220>
 <221> VARIANT
 <222> (11)..(12)
 <223> Xaa is any amino acid except Cys
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<220>
<221> VARIANT
<222> (14)..(16)
<223> Xaa is any amino acid except Cys
<400> 17
Asp Trp Val Cys Glu Asn Lys Lys Asp Gln Xaa Xaa Cys Xaa Xaa Xaa
<210> 18
<211> 16
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: variable
      sublibrary sequence used in designing focused
      secondary library
<220>
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<222> (6)..(7)
<223> Xaa is any amino acid except Cys
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<223> Xaa is any amino acid except Cys
<220>
<221> VARIANT
<222> (15)
<223> Xaa is any amino acid except Cys
<400> 18
Asp Trp Val Cys Glu Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Xaa Leu
<210> 19
<211> 16
<212> PRT
<213> Artificial Sequence
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      sublibrary sequence used in designing focused
      secondary library
<220>
<221> VARIANT
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<222> (5)..(7)
<223> Xaa is any amino acid except Cys
<220>
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<220>
<221> VARIANT
<222> (12)
<223> Xaa is any amino acid except Cys
<400> 19
Asn Trp Val Cys Xaa Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Ser Tyr
                                      10
<210> 20
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: variable
      sublibrary sequence used in designing focused
      secondary library
<220>
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<222> (1)
<223> Xaa is any amino acid except Cys
<220>
<221> VARIANT
<222> (3)
<223> Xaa is any amino acid except Cys
<220>
<221> VARIANT
<222> (14)..(16)
<223> Xaa is any amino acid except Cys
<400> 20
Xaa Trp Xaa Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Xaa Xaa Xaa
<210> 21
<211> 16
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: isolate of
      TN10/9 library found not to bind CEA
```

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<400> 21
Asn Trp Arg Cys Lys Leu Phe Pro Arg Tyr Pro Tyr Cys Ser Ser Trp
                                      10
<210> 22
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: isolate of
      TN10/9 library found not to bind CEA
<400> 22
Arg Tyr Cys Glu Phe Phe Pro Trp Ser Leu His Cys Gly Arg Pro
<210> 23
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: conserved
      amino acid positions in first family of CEA
      binding peptides
<220>
<221> VARIANT
<222> (6)
<223> X is Asn, Leu, Met or Phe
<220>
<221> VARIANT
<222> (7)
<223> X is Asp, Gly, Ile, Lys, Phe or Thr
<220>
<221> VARIANT
<222> (9)
<223> X is Arg, Asn, Asp, Glu or Gly
<220>
<221> VARIANT
<222> (12)
<223> X is Ala, Gly, His, Phe, Thr or Val
<220>
<221> VARIANT
<222> (15)
<223> X is Arg, Leu, Pro or Ser
<400> 23
Asp Trp Val Cys Glu Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Xaa Leu
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<210> 24
<211> 27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic CEA
      binding peptide with C-terminal immobilization
      sequence
<400> 24
Ser Asn Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asn Ser
                  5
                                     10
Tyr Ala Pro Gly Gly Glu Gly Gly Ser Lys
             20
<210> 25
<211> 27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic CEA
      binding peptide with C-terminal immobilization
      sequence
Ser Asp Trp Val Cys Glu Asn Lys Lys Asp Gln Trp Thr Cys Asn Leu
                  5
Leu Ala Pro Gly Gly Glu Gly Gly Ser Lys
             20
<210> 26
<211> 27
<212> PRT
<213> Artificial Sequence
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<220>

<223> Description of Artificial Sequence: synthetic CEA binding peptide with C-terminal immobilization sequence

Ser Asn Trp Asp Cys Met Phe Gly Ala Glu Gly Trp Ala Cys Ser Pro 15

Trp Ala Pro Gly Gly Glu Gly Gly Ser Lys 20 25

<210> 27

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<211> 27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic CEA
      binding peptide with C-terminal immobilization
      sequence
<400> 27
Ser Asp Trp Val Cys Glu Leu Thr Thr Gly Gly Tyr Val Cys Gln Pro
Leu Ala Pro Gly Gly Glu Gly Gly Ser Lys
             20
<210> 28
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: C-terminal
      sequence for immobilizing peptides
<400> 28
Ala Pro Gly Gly Glu Gly Gly Ser Lys
                  5
<210> 29
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: template
      sequence for sublibrary used in construction of
      focused secondary display library
<220>
<221> VARIANT
<222> (1)..(3)
<223> X is any amino acid except Cys
<220>
<221> VARIANT
<222> (5)..(6)
<223> X is any amino acid except Cys
<400> 29
Xaa Xaa Cys Xaa Xaa Lys Lys Asp Gln Trp Thr Cys Asn Leu Leu
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                                     10
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<210> 30

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<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: template
      sequence for sublibrary used in construction of
      focused secondary display library
<220>
<221> VARIANT
<222> (5)..(9)
<223> X is any amino acid except Cys
<400> 30
Asp Trp Val Cys Xaa Xaa Xaa Xaa Kaa Gln Trp Thr Cys Asn Leu Leu
<210> 31
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: template
      sequence for sublibrary used in construction of
      focused secondary display library
<220>
<221> VARIANT
<222> (8)..(12)
<223> X is any amino acid except Cys
<400> 31
Asp Trp Val Cys Glu Asn Lys Xaa Xaa Xaa Xaa Cys Asn Leu Leu
<210> 32
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: template
      sequence for sublibrary used in construction of
      focused secondary display library
<220>
<221> VARIANT
<222> (11) .. (12)
<223> X is any amino acid except Cys
<220>
<221> VARIANT
<222> (14)..(16)
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<223> X is any amino acid except Cys
<400> 32
Asp Trp Val Cys Glu Asn Lys Lys Asp Gln Xaa Xaa Cys Xaa Xaa
                                      10
<210> 33
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: template
      sequence for sublibrary used in construction of
      focused secondary display library
<220>
<221> VARIANT
<222> (6)..(7)
<223> X is any amino acid except Cys
<220>
<221> VARIANT
<222> (9)
<223> X is any amino acid except Cys
<220>
<221> VARIANT
<222> (12)
<223> X is any amino acid except Cys
<220>
<221> VARIANT
<222> (15)
<223> X is any amino acid except Cys
Asp Trp Val Cys Glu Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Xaa Leu
                  5
<210> 34
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: template
     sequence for sublibrary used in construction of
     focused secondary display library
<220>
<221> VARIANT
<222> (5)..(7)
<223> X is any amino acid except Cys
```

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<220>
<221> VARIANT
<222> (9)
<223> X is any amino acid except Cys
<220>
<221> VARIANT
<222> (12)
<223> X is any amino acid except Cys
<400> 34
Asn Trp Val Cys Xaa Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Ser Tyr
                  5
<210> 35
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: template
      sequence for sublibrary used in construction of
      focused secondary display library
<220>
<221> VARIANT
<222> (1)
<223> X is any amino acid except Cys
<220>
<221> VARIANT
<222> (3)
<223> X is any amino acid except Cys
<220>
<221> VARIANT
<222> (14)..(16)
<223> X is any amino acid except Cys
<400> 35
Xaa Trp Xaa Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Xaa Xaa Xaa
                                      10
<210> 36
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: family of CEA
      binding polypeptides
<220>
<221> VARIANT
<222> (1)
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<223> Xaa is Asp, Asn, Ala or Ile
<220>
<221> VARIANT
<222> (3)
<223> Xaa is Val, Ile, Met, Tyr, Phe, Pro or Asp
<220>
<221> VARIANT
<222> (5)
<223> Xaa is Asn, Glu or Asp
<220>
<221> VARIANT
<222> (6)
<223> Xaa is Leu, Phe, Typ, Trp, Val Met, Ile or Asn
<220>
<221> VARIANT
<222>.(7)
<223> Xaa is Phe, Leu, Asp, Glu, Ala, Ile, Lys, Asn,
     Ser, Val, Trp or Tyr
<220>
<221> VARIANT
<222> (8)
<223> Xaa is Lys, Phe, Asp, Gly, Leu, Asn or Trp
<220>
<221> VARIANT
<222> (9)
<223> Xaa is Asn, Pro, Phe, Gly, Asp, Ala, Ser, Glu, Gln
     or Trp
<220>
<221> VARIANT
<222> (10)
<223> Xaa is Gln or Lys
<220>
<221> VARIANT
<222> (12)
<223> Xaa is Phe, Thr, Met, Ser, Ala, Asn, Val, His,
      Ile, Pro, Trp or Tyr
<220>
<221> VARIANT
<222> (14)
<223> Xaa is Asn, Asp, Glu, Pro, Gln or Ser
<220>
<221> VARIANT
<222> (15)
<223> Xaa is Val, Leu, Ile, Pro, Ala, Gln, Ser, Met,
      Glu, Thr, Lys or Trp
<220>
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<221> VARIANT
<222> (16)
<223> Xaa is Leu, Met, Val, Tyr, Ala, Ile, Trp, His,
      Pro, Gln, Glu, Phe, Lys or Arg
<400> 36
Xaa Trp Xaa Cys Xaa Xaa Xaa Xaa Xaa Trp Xaa Cys Xaa Xaa Xaa
<210> 37
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
Asp Trp Met Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Leu Met
                  5
                                      10
                                                          15
<210> 38
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 38
Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Leu Met
                  5
                                      10
<210> 39
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 39
Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Gln Met
<210> 40
<211> 16
<212> PRT
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<213> Artificial Sequence

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<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 40
Asn Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Gln Glu
<210> 41
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 41
Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Gln Val Lys
                                      10
<210> 42
<211> 16
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 42
Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Met
 1
                  5
                                      10
<210> 43
<211> 16
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 43
Asp Trp Met Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Gln Ile
                  5
                                     10
                                                          15
<210> 44
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
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<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 44
Ile Trp Asp Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Pro Ala Pro
                  5
<210> 45
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Ile Arg
                  5
<210> 46
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 46
Asp Trp Met Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Val
                  5
                                      10
<210> 47
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 47
Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Ala Ile
<210> 48
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
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<223> Description of Artificial Sequence: CEA binding

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polypeptide
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<400> 48
Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Met Ala
                  5
                                      10
<210> 49
<211> 16
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 49
Asp Trp Val Cys Glu Phe Leu Lys Met Gln Trp Ala Cys Asn Val Leu
<210> 50
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asn Val Met
                  5
                                      10
<210> 51
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 51
Ala Trp Pro Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Pro Pro Gln
                  5
                                      10
 1
<210> 52
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
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<400> 52
Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Leu
                  5
 1
<210> 53
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 53
Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Lys Trp
                                      10
                  5
<210> 54
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 54
Asp Trp Val Cys Glu Trp Leu Lys Met Gln Trp Ala Cys Asn Met Leu
                                      10
                  5
<210> 55
<211> 16
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 55
Asp Trp Val Cys Asp Phe Phe Phe Asn Gln Trp Thr Cys Asn Leu Leu
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<223> Description of Artificial Sequence: CEA binding

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polypeptide
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Pro, Trp, Tyr, Gly, Leu or Glu
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Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys
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Trp, His, Arg, Met, Val, or Leu
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## 16-mer microprotein analogues

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16-mer microprotein analogues

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A DOCPHOENIX

APPL PARTS	NPL	CTNF
	Non-Patent Literature	Count Non-Final
IMIS	OATH	CTRS
Internal Misc. Paper	Oath or Declaration	Count Restriction
LET <u>V</u>	PET	EXIN
Misc. Incoming Letter	Petition	Examiner Interview
371P PCT Papers in a 371Application	RETMAIL Mail Returned by USPS	M903 DO/EO Acceptance
<u>.</u>	•	M905
AAmendment Including Elections	SEQLIST	DO/EO Missing Requirement
ABST	SPEC	NFDR
Abstract Abstract	Specification	Formal Drawing Required
ADS	SPEC NO	NOA
Application Data Sheet	Specification Not in English	Notice of Allowance
AF/D	TRNA	PETDEC
Affidavit or Exhibit Received	Transmittal New Application	Petition Decision
APPENDIX		
Appendix		
ARTIFACT	OUTGOING	INCOMING
Artifact	paners and the second s	
BIB	CTMS	AP.B
Bib Data Sheet	Misc. Office Action	Appeal Brief
CLM	1449 Signed 1449	C.AD Change of Address
	•	N/AP
COMPUTER Computer Program Listing	892	Notice of Appeal
CRFL	ABN	PA
All CRF Papers for Backfile	Abandonment	Change in Power of Attorney
DIST	APDEC	REM
Terminal Disclaimer Filed	Board of Appeals Decision	Applicant Remarks in Amendment
DRW	APEA	XT/
Drawings	Examiner Answer	Extension of Time filed separate
FOR	CTAV	
Foreign Reference	Count Advisory Action	
FRPR	CTEQ	
Foreign Priority Papers	Count Ex parte Quayle	
IDS	CTFR	File Wrapper
IDS Including 1449	Count Final Rejection	1
B4	ECBOX	FWCLM
Internal	Evidence Copy Box Identification	File Wrapper Claim
SRNT	WCLM	IIFW
Examiner Search Notes	Claim Worksheet	File Wrapper Issue Information
CLMPTO	WFEE	SRFW
PTO Prepared Complete Claim Set	Fee Worksheet	File Wrapper Search Info